

1547759_1.TXT

SEQUENCE LISTING

<110> Klein, Michael Schultz, Dennis

<120> Gene Mutation Associated with Age-Related Macular Degeneration

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Page 2

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570

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gaa Glu 234	Ile	ctg Leu	gat Asp	gaa Glu	ggt Gly 234	cac His 5	atc Ile	ctt Leu	cag Gln	ctg Leu 235	Lys	aac Asn	att Ile	cat His	gta Val 2355	7294
tct Ser	gac Asp	aca Thr	ggc Gly	cgt Arg 236	Tyr	gtg Val	tgt Cys	gtt Val	gct Ala 236	val	aat Asn	gta Val	gca Ala	gga Gly 2370	Met	7342
act Thr	gac Asp	aaa Lys	aaa Lys 237	Tyr	gac Asp	tta Leu	agt Ser	gtc Val 2380	His	gct Ala	cct Pro	cca Pro	agc Ser 238	Ile	ata Ile	7390
gga Gly	aac Asn	cac His 239	Arg	tca Ser	cct Pro	gaa Glu	aat Asn 239	Ile	agt Ser	gtg Val	gta Val	gaa Glu 2400	Lys	aac Asn	tca Ser	7438
gta Val	tct Ser 240	Leu	act Thr	tgt Cys	gaa Glu	gct Ala 2410	Ser	gga Gly	att Ile	CCC Pro	ctg Leu 241	Pro	tcc Ser	ata Ile	acc Thr	7486
tgg Trp 242	Phe	aaa Lys	gat Asp	ggg Gly	tgg Trp 242	cct Pro 5	gtc val	agc Ser	ctt Leu	agc Ser 2430	Asn	tct Ser	gtg Val	agg Arg	att Ile 2435	7534
ctt Leu	tca Ser	gga Gly	ggc Gly	agg Arg 2440	Met	cta Leu	cgg Arg	ctg Leu	atg Met 244	Gln	acc Thr	aca Thr	atg Met	gaa Glu 2450	Āsp	7582
gct Ala	ggc Gly	caa Gln	tat Tyr 245!	Thr	tgc Cys	gtt val	gta Val	agg Arg 2460	Asn	gca Ala	gct Ala	ggt Gly	gaa Glu 246!	Glu	aga Arg	7630
aaa Lys	atc Ile	ttt Phe 2470	Gly	ctt Leu	tca Ser	gta Val	tta Leu 247	Val	cca Pro	cct Pro	cat His	att Ile 2480	⊽a1	ggt Gly	gaa Glu	7678
aat Asn	aca Thr 248	Leu	gaa Glu	gat Asp	gtg Val	aag Lys 2490	Val	aaa Lys	gag Glu	aaa Lys	cag Gln 2499	Ser	gtt Val	acg Thr	ctg Leu	7726
act Thr 250	Cys	gaa Glu	gtg Val	aca Thr	ggg Gly 250!	aat Asn	cca Pro	gtg Val	cca Pro	gaa Glu 2510	Ile	aca Thr	tgg Trp	cac His	aaa Lys 2515	7774
gat Asp	ggg Gly	cag Gln	ccc Pro	ctc Leu 2520	Gln	gaa Glu	gat Asp	gaa Glu	gcc Ala 2525	His	cac His	att Ile	ata Ile	tct Ser 2530	Gly	7822
ggc Gly	cgt Arg	ttt Phe	ctt Leu 2535	Gin	att Ile	acc Thr	aat Asn	gtc Val 2540	Gln	gtg Val	cca Pro	cac His	act Thr 2545	Gly	aga Arg	7870
tat Tyr	aca Thr	tgt Cys 2550	Leu	gct Ala	tcc Ser	agt Ser	cca Pro 2555	Ala	ggc Gly	cac His	aag Lys	agc Ser 2560	Arg	agc Ser	ttc Phe	7918
agt Ser	ctt Leu 2565	Asn	gta Val	ttt Phe	gta Val	tct Ser 2570	Pro	aca Thr	att Ile	gct Ala	ggt Gly 2575	val	ggt Gly	agt Ser	gat Asp	7966
ggc Gly	aac Asn	cct Pro	gaa Glu	gat Asp	gtc Val	act Thr	gtc Val	atc Ile	Leu	aac Asn age	Ser	cct Pro	aca Thr	tct Ser	ttg Leu	8014

2380	2363	2390	2393
Val Cys Ğlu Ala 1		cca gct acc atc acc Pro Ala Thr Ile Thr 2605	
gat ggc act cct t Asp Gly Thr Pro 1 2615	ta gaa tct aac eu Glu Ser Asn	cga aat att cgt att Arg Asn Ile Arg Ile 2620	ctt cca gga 8110 Leu Pro Gly 2625
ggc aga act ctg of Gly Arg Thr Leu of 2630	ag atc ctc aat ln Ile Leu Asn 2635	gca cag gag gac aat Ala Gln Glu Asp Asn 264	Āla Ğly Arg
tac tct tgt gta g Tyr Ser Cys Val A 2645	cc acg aat gag la Thr Asn Glu 2650	gct gga gaa atg ata Ala Gly Glu Met Ile 2655	aag cac tat 8206 Lys His Tyr
gaa gtg aag gtg t Glu Val Lys Val 1 2660	ac att cca ccc yr Ile Pro Pro 2665	ata atc aat aaa ggg Ile Ile Asn Lys Gly 2670	gac ctt tgg 8254 Asp Leu Trp 2675
Gly Pro Gly Leu S	cc cct aaa gaa er Pro Lys Glu 680	gtg aag atc aaa gta Val Lys Ile Lys Val 2685	aac aac act 8302 Asn Asn Thr 2690
ctg acc ttg gaa t Leu Thr Leu Glu (2695	gt gaa gcg tat ys Glu Ala Tyr	gca att cct tct gcc Ala Ile Pro Ser Ala 2700	tcc ctc agc 8350 Ser Leu Ser 2705
tgg tac aag gat g Trp Tyr Lys Asp (2710	ga cag ccc ctt ly Gln Pro Leu 2715	aaa tcc gat gat cat Lys Ser Asp Asp His 272	Val Asn Ile
gct gcg aat gga d Ala Ala Asn Gly H 2725	ac aca ctt caa is Thr Leu Gln 2730	ata aag gag gct caa Ile Lys Glu Ala Gln 2735	ata tca gac 8446 Ile Ser Asp
acc gga cga tat a Thr Gly Arg Tyr 1 2740	ct tgt gta gca hr Cys Val Ala 2745	tct aac att gca ggt Ser Asn Ile Ala Gly 2750	gaa gat gag 8494 Glu Asp Glu 2755
Leu Asp Phe Asp V		gtt cct cca agt ttt Val Pro Pro Ser Phe 2765	
tgg gaa ata gga a Trp Glu Ile Gly A 2775	ac atg cta gat sn Met Leu Asp	act ggc agg aat ggt Thr Gly Arg Asn Gly 2780	gaa gcc aaa 8590 Glu Ala Lys 2785
gat gtg atc atc a Asp Val Ile Ile A 2790	ac aat ccc att sn Asn Pro Ile 2795	tct ctt tac tgt gag Ser Leu Tyr Cys Glu 280	Thr Asn Āla
gct ccc cct cct a Ala Pro Pro Pro T 2805	ca ctg aca tgg hr Leu Thr Trp 2810	tac aaa gat ggc cac Tyr Lys Asp Gly His 2815	cct ctg acc 8686 Pro Leu Thr
tca agt gat aaa g Ser Ser Asp Lys V 2820	ta ttg att ttg al Leu Ile Leu 2825	cca gga ggg cga gtg Pro Gly Gly Arg Val 2830	ttg cag att 8734 Leu Gln Ile 2835
cct cgg gct aaa g	ta gaa gat gct	ggg aga tac aca tgt Page 12	gtg gct gtg 8782

1547759_1.txt Pro Arg Ala Lys Val Glu Asp Ala Gly Arg Tyr Thr Cys Val Ala Val 2840 2845 2850 aat gag gct gga gaa gat tcc ctt caa tat gat gtc cgt gta ctc gtg Asn Glu Ala Gly Glu Asp Ser Leu Gln Tyr Asp Val Arg Val Leu Val 2855 2860 2865 ccg cca att atc aag gga gca aat agt gat ctc cct gaa gag gtc acc Pro Pro Ile Ile Lys Gly Ala Asn Ser Asp Leu Pro Glu Glu Val Thr gtg ctg gtg aac aag agt gca ctg ata gag tgt tta tcc agt ggc agc Val Leu Val Asn Lys Ser Ala Leu Ile Glu Cys Leu Ser Ser Gly Ser cca gca cca agg aat tcc tgg cag aaa gat gga cag ccc ttg cta gaa Pro Ala Pro Arg Asn Ser Trp Gln Lys Asp Gly Gln Pro Leu Leu Glu gat gac cat cat aaa ttt cta tct aat gga cga att ctg cag att ctg Asp Asp His His Lys Phe Leu Ser Asn Gly Arg Ile Leu Gln Ile Leu aat act caa ata aca gat atc ggc agg tat gtg tgt gct gag aac Asn Thr Gln Ile Thr Asp Ile Gly Arg Tyr Val Cys Val Ala Glu Asn aca gct ggg agt gcc aaa aaa tat ttt aac ctc aat gtt cat gtt cct Thr Ala Gly Ser Ala Lys Lys Tyr Phe Asn Leu Asn Val His Val Pro cca agt gtc att ggt cct aaa tct gaa aat ctt acc gtc gtg gtg aac Pro Ser Val Ile Gly Pro Lys Ser Glu Asn Leu Thr Val Val Val Asn aat ttc atc tct ttg acc tgt gag gtc tct ggt ttt cca cct cct gac Asn Phe Ile Ser Leu Thr Cys Glu Val Ser Gly Phe Pro Pro Pro Asp ctc agc tgg ctc aag aat gaa cag ccc atc aaa ctg aac aca aat act Leu Ser Trp Leu Lys Asn Glu Gln Pro Ile Lys Leu Asn Thr Asn Thr ctc att gtg cct ggt ggt cga act cta cag att att cgg gcc aag gta Leu Ile Val Pro Gly Gly Arg Thr Leu Gln Ile Ile Arg Ala Lys Val tca gat ggt ggt gaa tac act tgt ata gct atc aat caa gct ggc gaa Ser Asp Gly Glu Tyr Thr Cys Ile Ala Ile Asn Gln Ala Gly Glu agc aag aaa aag ttt tcc ctg act gtt tat gtg ccc cca agc att aaa Ser Lys Lys Phe Ser Leu Thr Val Tyr Val Pro Pro Ser Ile Lys gac cat gac agt gaa tct ctt tct gta gtt aat gta aga gag gga act Asp His Asp Ser Glu Ser Leu Ser Val Val Asp Val Arg Glu Gly Thr tct gtg tct ttg gag tgt gag tcg aac gct gtg cca cct cca gtc atc Ser Val Ser Leu Glu Cys Glu Ser Asn Ala Val Pro Pro Pro Val Ile

1547759_1.txt act tgg tat aag aat ggg cgg atg ata aca gag tct act cat gtg gag Thr Trp Tyr Lys Asn Gly Arg Met Ile Thr Glu Ser Thr His Val Glu att tta gct gat gga caa atg cta cac att aag aaa gct gag gta tct Ile Leu Ala Asp Gly Gln Met Leu His Ile Lys Lys Ala Glu Val Ser gac aca ggc cag tat gta tgt aga gct ata aat gta gca gga cgg gat ÁSP Thr Gly Gln Tyr Val Cýs Arg Ála Ile Asn Val Ála Gly Arg ÁSP gat aaa aat ttc cac ctc aat gta tat gtg cca ccc agt att gaa gga Ăsp Lys Asn Phe His Leu Asn Val Tyr Val Pro Pro Ser Ile Ğlu Ğly cct gaa aga gaa gtg att gtg gag acg atc agc aat cct gtg aca tta Pro Glu Arg Glu Val Ile Val Glu Thr Ile Ser Asn Pro Val Thr Leu aca tgt gat gcc act ggg atc cca cct ccc acg ata gca tgg tta aag Thr Cys Asp Ala Thr Gly Ile Pro Pro Pro Thr Ile Ala Trp Leu Lys aac cac aag cgc ata gaa aat tct gac tca ctg gaa gtt cgt att ttg Asn His Lys Arg Ile Ğlu Asn Ser Asp Ser Leŭ Ğlu Val Arg Ile Leŭ tct gga ggt agc aaa ctc cag att gcc cgg tct cag cat tca gat agt Ser Gly Gly Ser Lys Leu Gln Ile Ala Arg Ser Gln His Ser Asp Ser gga aac tat aca tgt att gct tca aat atg gag gga aaa gcc cag aaa Gly Asn Tyr Thr Cys Ile Ala Ser Asn Met Glu Gly Lys Ala Gln Lys tat tac ttt ctt tca att caa gtt cct cca agt gtt gct ggt gct gaa Tyr Tyr Phe Leu Ser Ile Gln Val Pro Pro Ser Val Ala Gly Ala Glu att cca agt gat gtc agt gtc ctt cta gga gaa aat gtt gag ctg gtc Ile Pro Ser Asp Val Ser Val Leu Leu Gly Glu Asn Val Glu Leu Val tgc aat gca aat ggc att cct act cca ctt att caa tgg ctt aaa gat Cys Asn Ala Asn Gly Ile Pro Thr Pro Leu Ile Gln Trp Leu Lys Asp gga aag ccc ata gct agt ggt gaa aca gaa aga atc cga gtg agt gca Gly Lys Pro Ile Ala Ser Gly Glu Thr Glu Arg Ile Arg Val Ser Ala aat ggc agc aca tta aac att tat gga gct ctt aca tct gac acg ggg Asn Gly Ser Thr Leu Asn Ile Tyr Gly Ala Leu Thr Ser Asp Thr Gly 3300 3305 3310 331 aaa tac aca tgt gtt gct act aat ccc gct gga gaa gaa gac cga att Lys Tyr Thr Cys Val Ala Thr Asn Pro Ala Gly Glu Glu Asp Arg Ile ttt aac ttg aat gtc tat gtt aca cct aca att agg ggt aat aaa gat Phe Asn Leu Asn Val Tyr Val Thr Pro Thr Ile Arg Gly Asn Lys Asp

Page 14

gaa gca gag aaa Glu Ala Glu Lys 3350	cta atg act Leu Met Thr	tta gtg gat Leu Val Asp 3355	act tca ata Thr Ser Ile 336	Asn Ile Glu	10318
tgc aga gcc aca Cys Arg Ala Thr 3365	ggg acg cct Gly Thr Pro 337	Pro Pro Gln	ata aac tgg Ile Asn Trp 3375	ctg aag aat Leu Lys Asn	10366
gga ctt cct ctg Gly Leu Pro Leu 3380					10414
caa gtt atc agg Gln Val Ile Arg	att gtg aga Ile Val Arg 3400	gct cag gtg Ala Gln Val 340	Ser Asp Val	gct gtg tat Ala Val Tyr 3410	10462
act tgt gtg gcc Thr Cys Val Ala 341		gct ggg gtg Ala Gly Val 3420	gat aat aag Asp Asn Lys	cat tac aat His Tyr Asn 3425	10510
ctt caa gtg ttt Leu Gln val Phe 3430				Ğİÿ Thr Ğlü	10558
gaa atc aca gtt Glu Ile Thr Val 3445	ctc aaa ggt Leu Lys Gly 345	Ser Ser Thr	tct atg gca Ser Met Ala 3455	tgc att act Cys Ile Thr	10606
gat gga acc cca Asp Gly Thr Pro 3460	gct ccc agt Ala Pro Ser 3465	atg gcc tgg Met Ala Trp	ctt aga gat Leu Arg Asp 3470	ggc cag cct Gly Gln Pro 3475	10654
ctg ggg ctt gat Leu Gly Leu Asp	gcc cat ctg Ala His Leu 3480	aca gtc agc Thr Val Ser 348	Thr His Gly	atg gtc ctg Met Val Leu 3490	10702
cag ctc ctc aaa Gln Leu Leu Lys 349	Ala Glu Thr	gaa gat tcg Glu Asp Ser 3500	gga aag tac Gly Lys Tyr	acc tgc att Thr Cys Ile 3505	10750
gcc tca aat gaa Ala Ser Asn Glu 3510	gct gga gaa Ala Gly Glu	gtc agc aag Val Ser Lys 3515	cac ttt atc His Phe Ile 352	Leu Lys Val	10798
cta gaa cca cct Leu Glu Pro Pro 3525	cac att aat His Ile Asn 353	Gly Ser Glu	gaa cat gaa Glu His Glu 3535	gag ata tca Glu Ile Ser	10846
gta att gtt aat Val Ile Val Asn 3540	aac cca ctt Asn Pro Leu 3545	gaa ctt acc Glu Leu Thr	tgc att gct Cys Ile Ala 3550	tct gga atc Ser Gly Ile 3555	10894
cca gcc cct aaa Pro Ala Pro Lys			Gly Arg Pro		10942
acg gat caa gtg Thr Asp Gln Val 357	Gln Thr Leu	gga gga gga Gly Gly Gly 3580	gag gtt ctt Glu Val Leu	cga att tct Arg Ile Ser 3585	10990
act gct cag gtg Thr Ala Gln Val	gag gat aca	gga aga tat	aça tgt ctg	gca tcc agt	11038

cct gca gga Pro Ala Gly 3605	gat gat gat Asp Asp Asp	aag gaa tat Lys Glu Tyr 3610	cta gtg aga Leu Val Arg 361		a cct 11086 I Pro
cct aat att Pro Asn Ile 3620	gct gga act Ala Gly Thr 362	Asp Glu Pro	cgg gat atc Arg Asp Ile 3630	act gtg tta Thr Val Le	a cgg 11134 u Arg 3635
aac aga caa Asn Arg Gln	gtg aca ttg Val Thr Leu 3640	gaa tgc aag Glu Cys Lys	tca gat gca Ser Asp Ala 3645	gtg ccc ccc Val Pro Pro 36	Pro
gta att act Val Ile Thr	tgg ctc aga Trp Leu Arg 3655	aat gga gaa Asn Gly Glu 366	cgg tta cag Arg Leu Gln O	gca aca cca Ala Thr Pro 3665	t cga 11230 o Arg
gtg cga atc Val Arg Ile 3670	Leu Ser Gly	ggg aga tac Gly Arg Tyr 3675	ttg caa atc Leu Gln Ile	aac aat gc Asn Asn Ala 3680	t gac 11278 a Asp
cta ggt gat Leu Gly Asp 3685	aca gcc aat Thr Ala Asn	tat acc tgt Tyr Thr Cys 3690	gtt gcc agc Val Ala Ser 369	Asn Ile Ala	a gga 11326 a Gly
		_Ile Leu Thr	gta aat gtt Val Asn Val 3710		
aag ggg ggc Lys Gly Gly	ccc cag agc Pro Gln Ser 3720	ctt gta att Leu Val Ile	ctt tta aat Leu Leu Asn 3725	aag tca act Lys Ser Thi 37:	r val
ttg gaa tgc Leu Glu Cys	atc gct gaa Ile Ala Glu 3735	ggt gtg cca Gly Val Pro 374	act cca agg Thr Pro Arg O	ata aca tgg Ile Thr Trp 3745	g aga 11470 o Arg
aag gat gga Lys Asp Gly 3750	Ala Val Leu	gct ggg aat Ala Gly Asn 3755	cat gca aga His Ala Arg	tat tcc atc Tyr Ser Ile 3760	ttg 11518 Leu
			gca cat gtc Ala His Val 377	Thr Asp Thi	
cgg tat ttg Arg Tyr Leu 3780	tgt atg gcc Cys Met Ala 378	_Thr Asn Ala	gct gga aca Ala Gly Thr 3790	gat cgc agg Asp Arg Arg	g cga 11614 g Arg 3795
ata gat tta Ile Asp Leu	cag gtc cat Gln Val His 3800	gtt cct cca Val Pro Pro	tct att gct Ser Ile Ala 3805	ccg ggt cct Pro Gly Pro 381	Thr
Asn Met Thr	gta ata gta Val Ile Val 3815	aat gtt caa Asn Val Gln 382	act act ctg Thr Thr Leu O	gct tgt gad Ala Cys Gli 3825	g gct 11710 i Ala
	Pro Lys Pro		tgg aga aaa Trp Arg Lys		
ctt aat gtg	gat caa aat	cag aac tca	tac agg ctc Page 16	ctt tct tca	a ggt 11806

Leu	Asn 3845		Asp	Gln	Asn	G]n 3850		Ser	1547 Tyr	759 <u> </u> Arg	1.tx Leu 3855	Leu	Ser	Ser	Gly	
tca Ser 3860	Leu	gta Val	att Ile	att Ile	tcc Ser 3865	cct Pro	tct Ser	gtg Val	gat Asp	gac Asp 3870	Thr	gca Ala	acc Thr	tat Tyr	gaa Glu 3875	11854
tgt Cys	act Thr	gtg Val	aca Thr	aac Asn 3880	Gly	gct Ala	gga Gly	gat Asp	gat Asp 3885	Lys	aga Arg	act Thr	gtg Val	gat Asp 3890	Leu	11902
act Thr	gtc Val	caa Gln	gtt Val 3899	Pro	cct Pro	tcc Ser	ata Ile	gct Ala 3900	Asp	gag Glu	cct Pro	aca Thr	gat Asp 3905	Phe	cta Leu	11950
gta Val	acc Thr	aaa Lys 3910	His	gcc Ala	cca Pro	gca Ala	gta Val 3919	Ile	acc Thr	tgc Cys	act Thr	gct Ala 3920	Ser	gga Gly	gtt Val	11998
		Pro				tgg Trp 393(Thr					Arg				12046
	Ğİy					att Ile					Āla					12094
gcc Ala	acc Thr	caa Gln	tta Leu	aac Asn 3960	His	gct Ala	gga Gly	aga Arg	tac Tyr 3969	Thr	tgt Cys	gtc Val	gct Ala	agg Arg 3970	Asn	12142
gcg Ala	gct Ala	ggc Gly	tct Ser 3975	Āla	cat His	cga Arg	cac His	gtg Val 3980	Thr	ctt Leu	cat His	gtt Val	cat His 3985	Glu	cct Pro	12190
cca Pro	gtc Val	att Ile 3990	Gln	ccc Pro	caa Gln	cca Pro	agt Ser 399	Ğlu	cta Leu	cac His	gtc Val	att Ile 4000	Leu	aac Asn	aat Asn	12238
cct Pro	att Ile 4005	Leu	tta Leu	cca Pro	tgt Cys	gaa Glu 401(Ala	aca Thr	ggg Gly	aca Thr	ccc Pro 4015	Ser	cct Pro	ttc Phe	att Ile	12286
	Trp					atc Ile					Ser					12334
gca Ala	gtt Val	ctt Leu	cct Pro	agt Ser 4040	Gly	ggc Gly	tta Leu	cag Gln	atc Ile 4045	Ser	aga Arg	gct Ala	gtc Val	cga Arg 4050	Glu	12382
gat Asp	gct Ala	ggc Gly	act Thr 4055	Tyr	atg Met	tgt Cys	gtg Val	gcc Ala 4060	Gln	aac Asn	ccg Pro	gct Ala	ggt Gly 4065	Thr	gcc Ala	12430
			Ile			aat Asn		Gln					Ile			12478
cat His	cta Leu 4085	Lys	gaa Glu	tat Tyr	gtt Val	att Ile 4090	Āla	gtg Val	gac Asp	aag Lys	ccc Pro 4095	Ile	acg Thr	tta Leu	tcc Ser	12526

1547759_1.txt ct gac att aca tgg ca

tgt gaa gca gat Cys Glu Ala Asp 4100	ggc ctc cct c Gly Leu Pro P 4105	ccg cct gac at Pro Pro Asp Il 41	t aca tgg cat e Thr Trp His	aaa gat 12574 Lys Asp 4115
ggg cgt gca att Gly Arg Ala Ile	gtg gaa tct a Val Glu Ser I 4120	itc cgc cag cg Tle Arg Gln Ar 4125	c gtc ctc agc g Val Leu Ser	tct ggc 12622 Ser Gly 4130
tct ctg caa ata Ser Leu Gln Ile 413	Ala Phe Val G	ag cct ggt ga In Pro Gly As 4140	t gct ggc cat p Ala Gly His 4145	Tyr Thr
tgc atg gca gcc Cys Met Ala Ala 4150	Asn Val Ala G	ga tca agc ag lly Ser Ser Se 155	c aca agc acc r Thr Ser Thr 4160	aag ctc 12718 Lys Leu
acc gtc cat gta Thr Val His Val 4165	cca ccc agg a Pro Pro Arg I 4170	itc aga agt ac Te Arg Ser Th	a gaa gga cac r Glu Gly His 4175	tac acg 12766 Tyr Thr
gtc aat gag aat Val Asn Glu Asn 4180			s Val Ala Asp	
ccc aca cca gca Pro Thr Pro Ala				
ttg tta gga aaa Leu Leu Gly Lys 421	${}_{ extsf{T}}$ Tyr Thr A ${}^{ extsf{T}}$ a G	gaa cca tat gg Glu Pro Tyr Gl 4220	a gaa ctc att y Glu Leu Ile 4225	Leu Glu
aat gtt gtg ctg Asn Val Val Leu 4230	Glu Asp Ser G	gc ttc tat ac ily Phe Tyr Th 1235	c tgt gtt gct r Cys Val Ala 4240	aac aat 12958 Asn Asn
gct gca ggt gaa Ala Ala Gly Glu 4245	gat aca cac a Asp Thr His T 4250	ct gtc agc ct hr Val Ser Le	g act gtg cat u Thr Val His 4255	gtt ctc 13006 Val Leu
ccc act ttt act Pro Thr Phe Thr 4260	gaa ctt cct g Glu Leu Pro G 4265	ga gac gtg tc ly Asp Val Se 42	r Leu Asn Lys	gga gaa 13054 Gly Glu 4275
cag cta cga tta Gln Leu Arg Leu	agc tgt aaa g Ser Cys Lys A 4280	ct act ggt at la Thr Gly Il 4285	t cca ttg ccc e Pro Leu Pro	aaa tta 13102 Lys Leu 4290
aca tgg acc ttc Thr Trp Thr Phe 429	Asn Asn Asn I	itt att cca gc Tle Ile Pro Al 4300	c cac ttt gac a His Phe Asp 4305	Ser Val
aat gga cac agt Asn Gly His Ser 4310	Ğlu Leu Val I	itt gaa aga gt le Glu Arg Va 315	g tca aaa gag 1 Ser Lys Glu 4320	gat tca 13198 Asp Ser
ggt act tat gtg Gly Thr Tyr Val 4325	tgc acc gca g Cys Thr Ala G 4330	ag aac agc gt ilu Asn Ser Va	t ggc ttt gtg 1 Gly Phe Val 4335	aag gca 13246 Lys Ala
att gga ttt gtt Ile Gly Phe Val 4340	tat gtg aaa g Tyr Val Lys G 4345	aa cct cca gt ilu Pro Pro Va 43 Page	l Phe Lys Gly 50	gat tat 13294 Asp Tyr 4355

cct tct aac Pro Ser Asn	tgg att gg Trp Ile G 4360	aa cca cti lu Pro Lei	ıĞlyĞl	gg aat gca ly Asn Ala 865	atc ctg Ile Leu	aat tgi Asn Cys 4370	13342
gag gtg aaa Glu Val Lys	gga gac co Gly Asp P 4375	cc acc cca ro Thr Pro	a acc at Thr Il 4380	c cag tgg le Gln Trp	aac aga Asn Arg 438	_Lys Gly	a 13390
gtg gat att Val Asp Ile 4390	Glu Ile Si	gc cac aga er His Arq 439	g Ile Ar	gg caa ctg g Gln Leu	ggc aat Gly Asn 4400	ggc tco Gly Sei	13438
ctg gcc atc Leu Ala Ile 4405	tat ggc a Tyr Gly Tl	et gtt aa ir Val Asi 4410	t gaa ga n Glu As	at gcc ggt sp Ala Gly 441	Asp Tyr	aca tgi Thr Cys	13486
gta gct acc Val Ala Thr 4420	aat gaa g Asn Glu A 44	ct ggg gtg la Gly Va 125	gtg ga I Val Gl	ng cgc agc lu Arg Ser 4430	atg agt Met Ser	ctg act Leu Thi 443	13534
ctg caa agt Leu Gln Ser	cct cct at Pro Pro I 4440	t atc ac le Ile Th	¹ Leu Ğl	ig cca gtg lu Pro Val 145	gaa act Glu Thr	gtt att Val Ile 4450	13582
aat gct ggt Asn Ala Gly	ggc aaa a Gly Lys I 4455	c ata tte le Ile Lei	aat to Asn Cy 4460	gt cag gca ⁄s Gln Ala	act gga Thr Gly 446	_Glu Pro	13630
caa cca acc Gln Pro Thr 447(Ile Thr Ti	gg tcc cg p Ser Are 44	g Gln Ğl	g cac tct y His Ser	att tcc Ile Ser 4480	tgg gat Trp Asp	13678
gac cgg gtt Asp Arg Val 4485	aac gtg ti Asn Val Le	g tcc aad eu Ser Asi 4490	aac to n Asn Se	ta tta tat er Leu Tyr 449	Ile Āla	gat gci Asp Ala	13726 a
cag aaa gaa Gln Lys Glu 4500	Asp Thr Se	t gaa tt er Glu Pho 505	t gaa to e Glu Cy	gt gtt gct vs Val Ala 4510	cga aac Arg Asn	tta ato Leu Met 451	:
ggt tct gtc Gly Ser Val	ctt gtc ag Leu Val Ai 4520	ga gtg cca rg Val Pro	o Val Il	a gtc cag le Val Gln i25	gtt cat Val His	ggt gga Gly Gly 4530	a 13822 /
ttt tcc cag Phe Ser Gln	tgg tct gg Trp Ser A 4535	a tgg aga la Trp Arg	g gcc to g Ala Cy 4540	gc agt gtc vs Ser Val	acc tgt Thr Cys 454	Gly Lys	13870 5
ggc atc caa Gly Ile Gln 4550	Lys Arg Se		i Cys As				
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16318

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Gly Ala Glu Ile Pro Ser Asp Val Ser Val Leu Leu Gly Glu Asn Val

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16,263 of SEQ ID NO:1

<400> 24 caggacaaca tttatt	16
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<210> <211> <212> <213>	> 17	
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	16,263 of SEQ ID NO:1	
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	1547750 1 + v+	
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16,263 of SEQ ID NO:1

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	134//39_1. LXL	
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21

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	16,263 of SEQ ID NO:1	
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	134//39_1. CXC	
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16,263 of SEQ ID NO:1

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16,263 of SEQ ID NO:1

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16,263 of SEQ ID NO:1

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1547759_1.txt
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Pro Pro Gly Gln His Leu Leu Gly Asp Gly Lys Ser Cys Ala
20 25 30
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